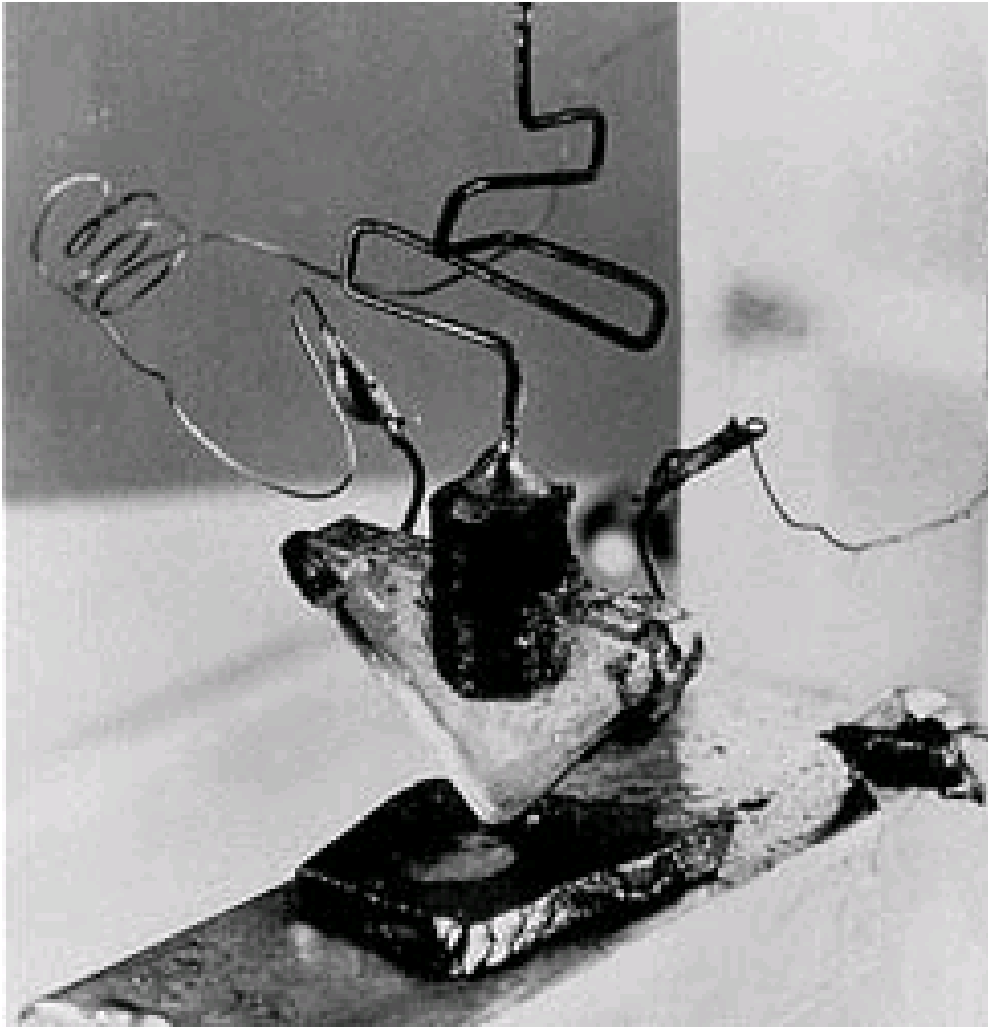




# *iGEM program overview*

*March 18, 2008*  
Andrew Hessel  
[ahessel@gmail.com](mailto:ahessel@gmail.com)

## Foundation of IT



1947

## Foundation of Bioengineering



1953

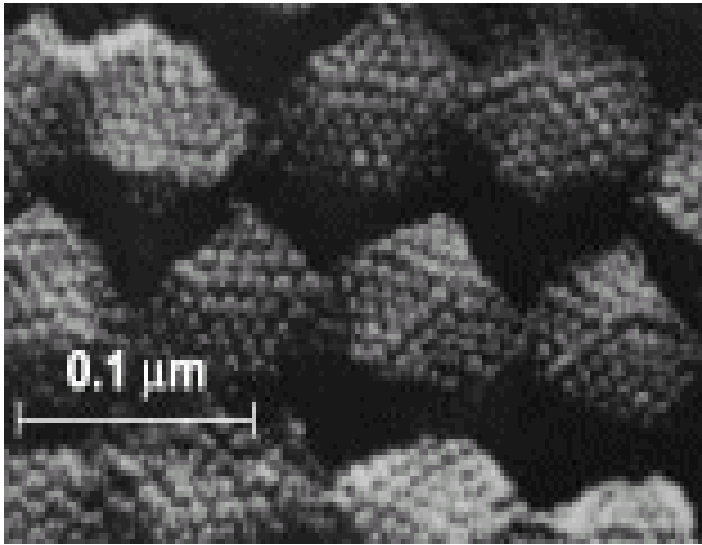




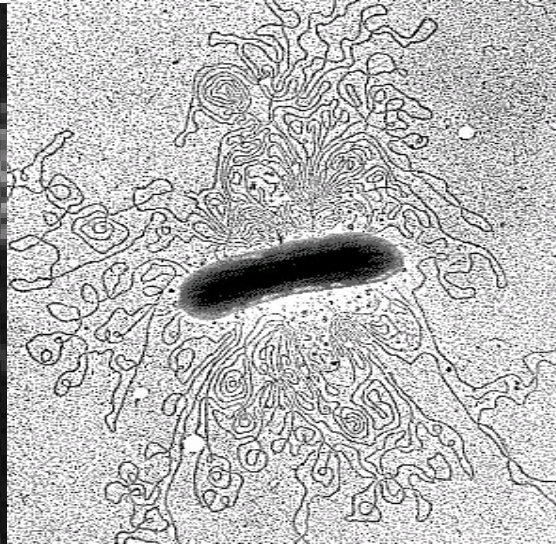
8000X for \$49.99







50 KB



5 MB



6 GB

DNA is a programming language for biochemistry (metabolism).

It can also specify how to build hardware.

Nature supports incredible diversity – over 2 million species we can see!

ATG-TCA-AAT-AAA-AGT-AAT-GAT-AAT-GGC-AGA-GCA-TAT-GAG-TTT-GCA-TTT-ATA-AAT-GAA-  
TTA-GGA-CGC-ATT-GCA-ACT-CAA-AAT-CAG-AAT-ATA-AAT-ATC-GAA-AAG-AAT-TCT-AGC-TAT-  
TAC-GTA-GTT-GAG-AAA-TCT-TGG-AGT-ACA-TTA-TCG-GAT-CTT-GAA-AAA-GAA-AAA-TAT-ACA-  
AAA-AGT-GCA-ATT-GCT-GGT-ATC-AAT-CTT-ATA-ACA-AGC-TTA-GAG-CCA-ATA-ATA-GAA-GAT-  
GGT-AAT-GGT-GTA-TTA-AAC-TTA-AAA-ATA-CAA-GCT-GAT-AAT-AAA-GGT-GAA-TTA-GGC-GAT-  
ATT-AGA-GAT-ATT-TTA-ATT-CAA-AGA-GAA-AAT-ATT-AAT-TGG-GAA-ATT-GGT-TTA-AGT-TTA-  
AAA-CAT-AAT-CAT-TTT-GCT-GTG-AAA-CAT-AGT-CGT-TTA-TCA-CAT-AAA-ATT-GAT-TTT-TCA-  
GAA-AAA-TGG-TTC-CAA-TTA-CCT-TCT-TCT-CAA-AAT-TAT-TGG-GAT-AAT-ATA-CTC-CCT-ATT-  
TTT-GAG-AAA-TTA-GAA-ATT-TAT-AAA-AAA-GAT-AAA-ATA-AAA-TGG-AGA-GAG-TTA-TCT-AAT-  
AAA-GAA-GAT-TGC-ATT-TAT-TAT-CCC-ATA-CTT-AAA-TCA-TTT-ATA-GCA-GAA-ATT-AAA-GAA-  
AAG-TAT-GAT-AAA-TAT-AAT-TCT-ATT-GTT-CCA-CAG-AGA-ATG-GTT-GAA-TAT-TTA-CTT-GGA-  
TAT-TTT-GAT-TTC-TAT-AAA-ATC-ATA-AGT-CAA-GAT-AAT-AAG-AAA-CTA-ACA-TCT-ATT-CAA-  
TCA-TTT-AAT-TTA-CGT-GGA-ACA-CTA-AAT-AAA-CCC-TCT-AAA-AAA-CGA-AAG-GCA-GAC-ATT-  
TTT-ATA-CCT-GTA-GCT-AAT-TTA-CCA-ACT-AGA-ATC-ATT-GAT-ATA-GAT-TTT-AAG-CCA-AAT-  
AGT-AAA-AAC-ACG-GTT-GAA-TTA-TAT-TTA-GAT-AAA-GGA-TGG-CAA-TTT-AGT-TTT-AGA-ATA-  
CAT-AAT-GCT-TCT-ACT-ATT-ATT-GAA-CCG-AGC-TTG-AAA-TTT-GAT-ATA-AAA-CTT-ATT-GGT-  
GTT-CCT-GCC-ACA-ATA-ATT-TGT-TTA-GAG-ACC-CCT-TGG-GAA-GAA-TGA



1980  
500 bp/day (manual)



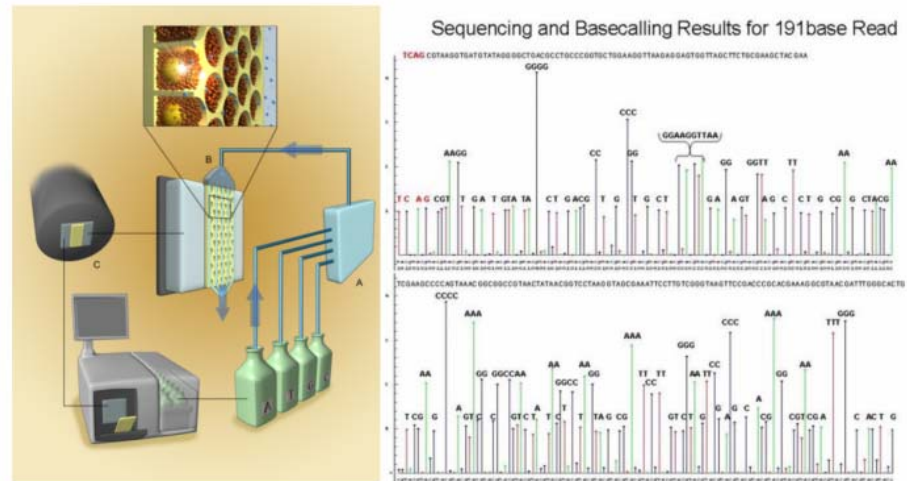
1987  
36,000 bp/day (semi-auto)



1995  
144,000 bp/day (semi-auto)



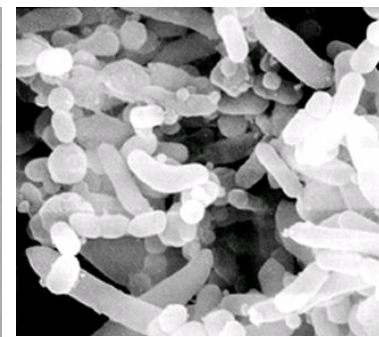
1998  
500,000 bp/day (automatic)



## 2007 – Sequencing by Synthesis

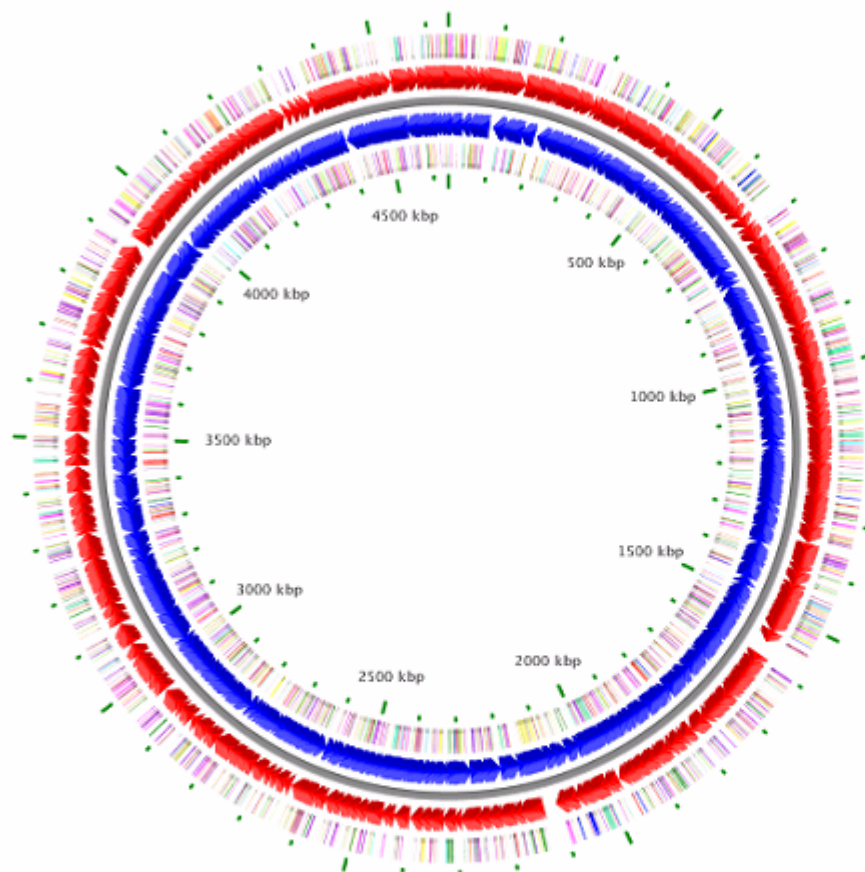


## Environmental Shotgun Sequencing: Its Potential and Challenges for Studying the Hidden World of Microbes





*E coli K12 complete genome*



# BASys

## Genes encoding proteins

- Forward strand
- Reverse strand

## Genes encoding functional RNA

- Forward strand
- Reverse strand

## COG functional categories

### Information storage and processing

- Translation, ribosomal structure and biogenesis
- Transcription
- DNA replication, recombination and repair

### Cellular processes

- Cell division and chromosome partitioning
- Posttranslational modification, protein turnover, chaperones
- Cell envelope biogenesis, outer membrane
- Cell motility and secretion
- Inorganic ion transport and metabolism
- Signal transduction mechanisms

### Metabolism

- Energy production and conversion
- Carbohydrate transport and metabolism
- Amino acid transport and metabolism
- Nucleotide transport and metabolism
- Coenzyme metabolism
- Lipid metabolism
- Secondary metabolites biosynthesis, transport and catabolism

### Poorly characterized

- General function prediction only
- Function unknown

BASys: Friday April 15 09:42:20 2005

Length: 4,639,675 bp; Genes: 4,254

Expand -

Expand +

Full view

Rotate -

Rotate +

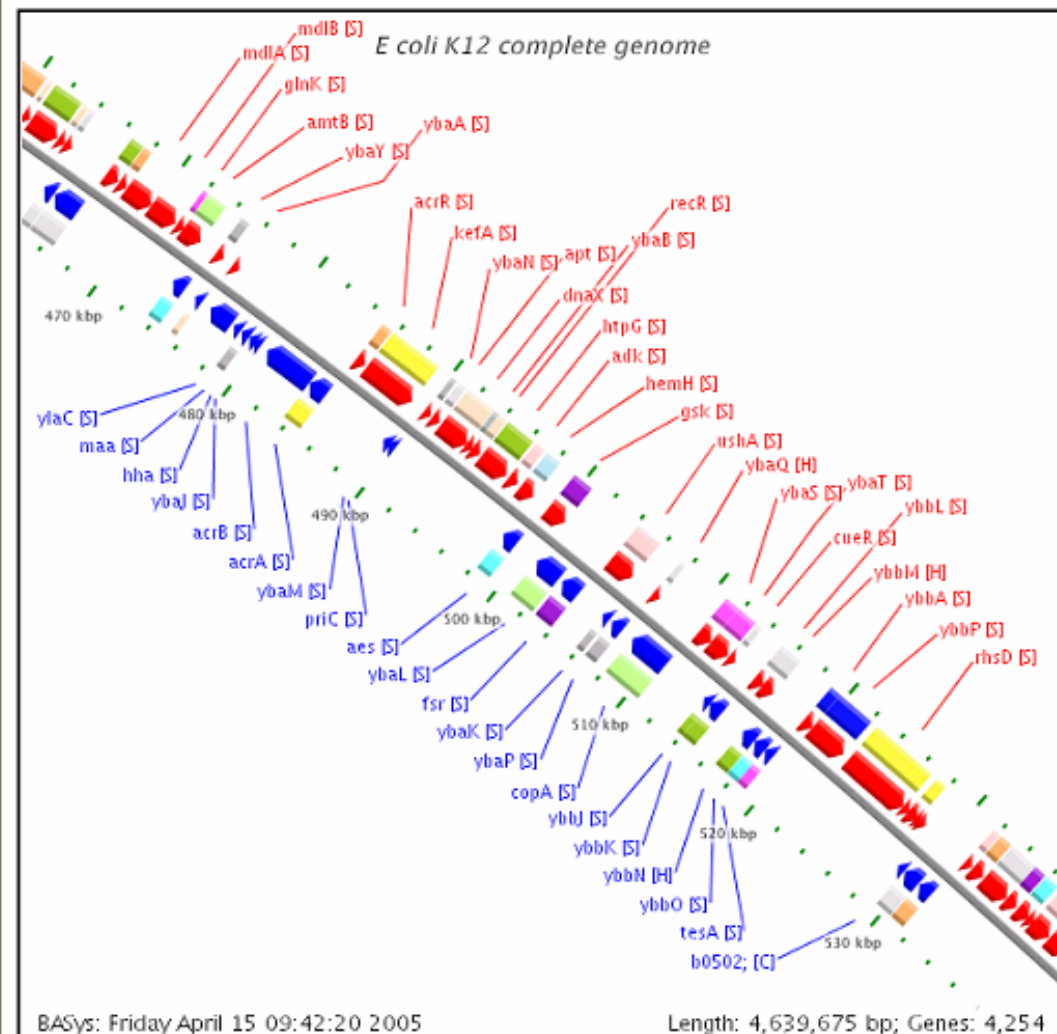
?

Click tick marks to expand the view.

Valid XHTML 1.0; Valid CSS.

Displayed PNG file size: 188 kb.

Centered on base 1; Zoom = 1.



# BASys

## Genes encoding proteins

- Forward strand
- Reverse strand

## Genes encoding functional RNA

- Forward strand
- Reverse strand

## COG functional categories

### Information storage and processing

- Translation, ribosomal structure and biogenesis
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- DNA replication, recombination and repair

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### Metabolism

- Energy production and conversion
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- Amino acid transport and metabolism
- Nucleotide transport and metabolism
- Coenzyme metabolism
- Lipid metabolism
- Secondary metabolites biosynthesis, transport and catabolism

### Poorly characterized

- General function prediction only
- Function unknown

Expand - Expand + Full view Rotate - Rotate + ?

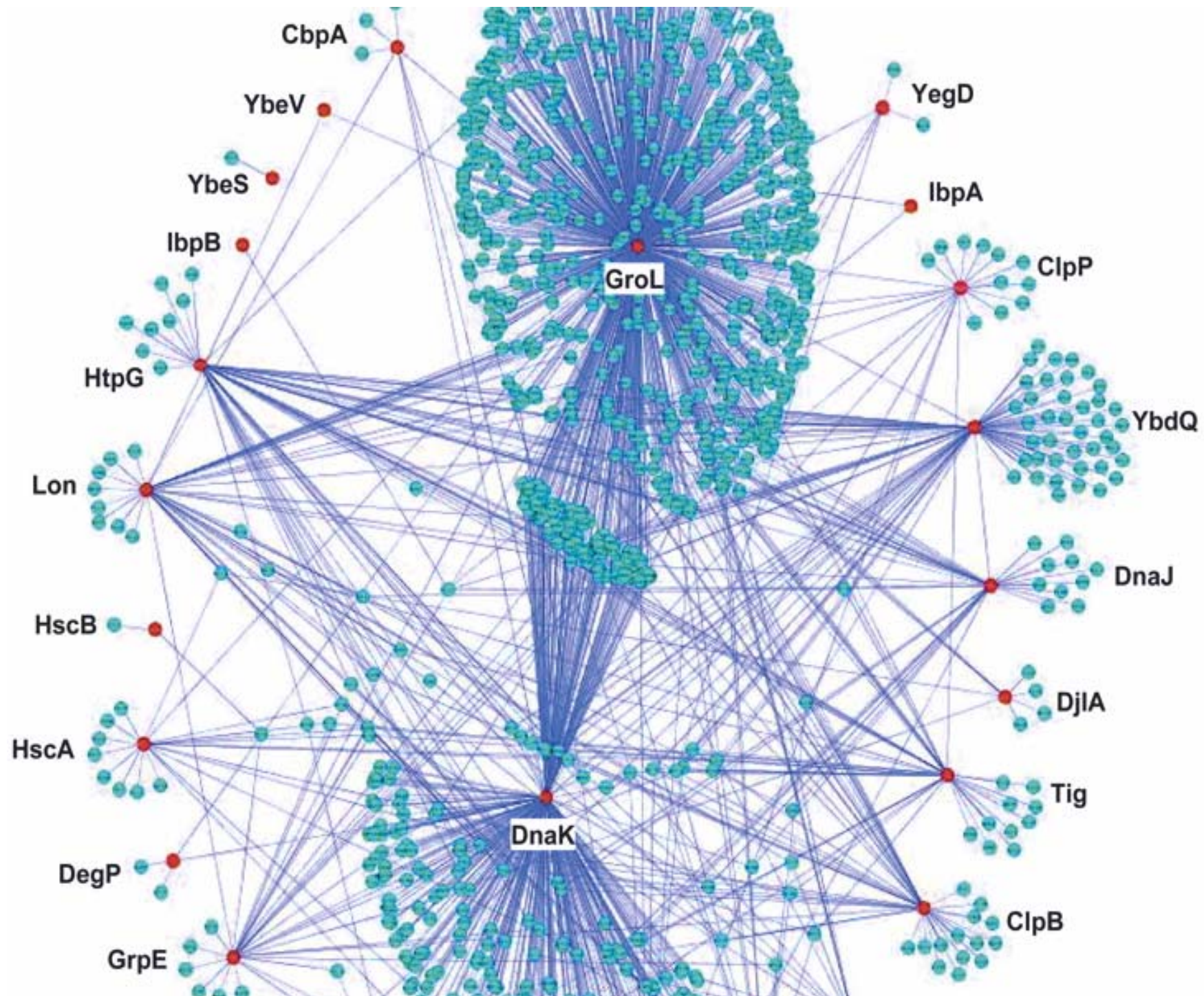
This is a fully expanded view.

Valid XHTML 1.0; Valid CSS.

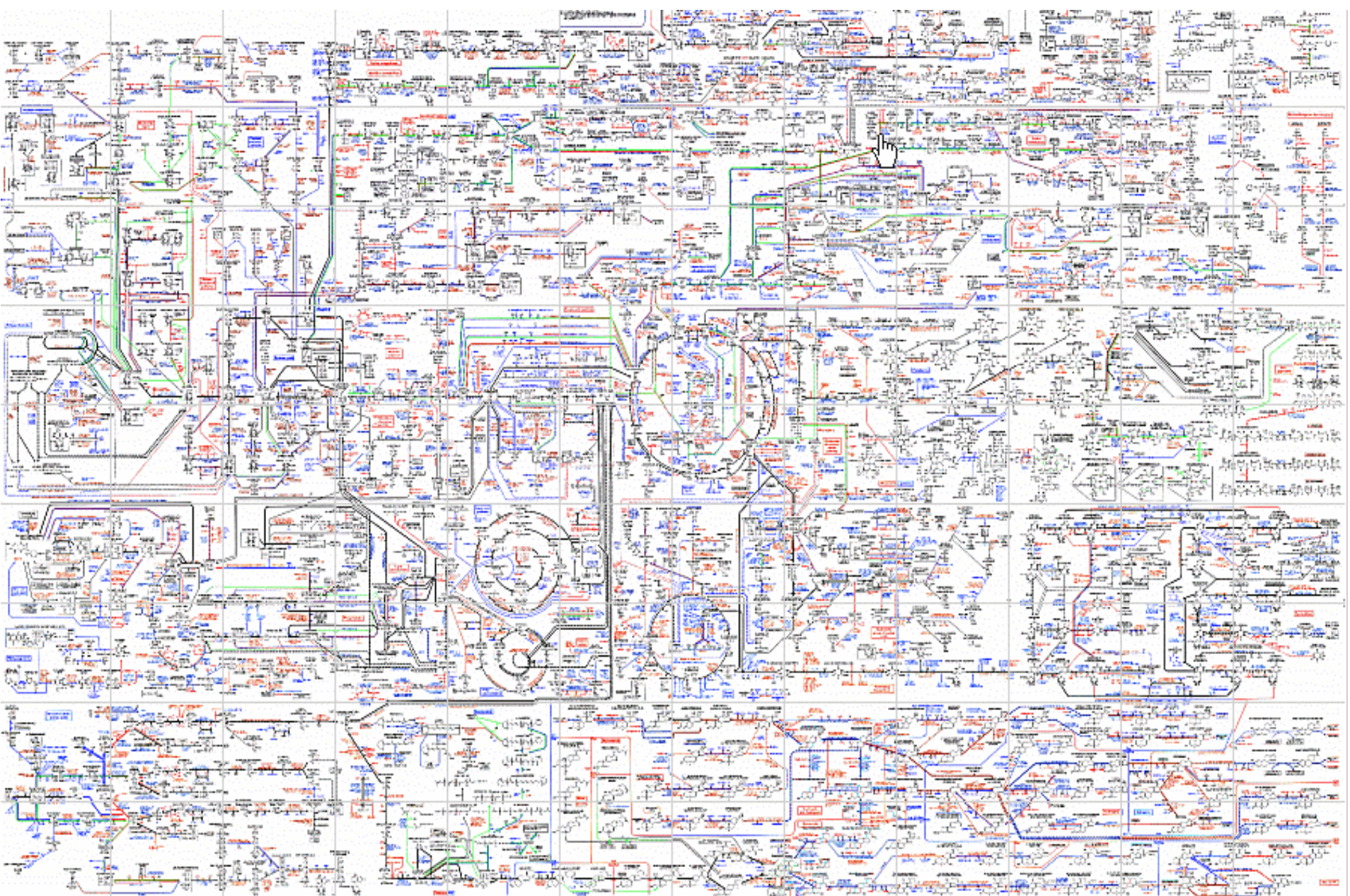
Displayed PNG file size: 79 kb.

Centered on base 500,000; Zoom = 36.







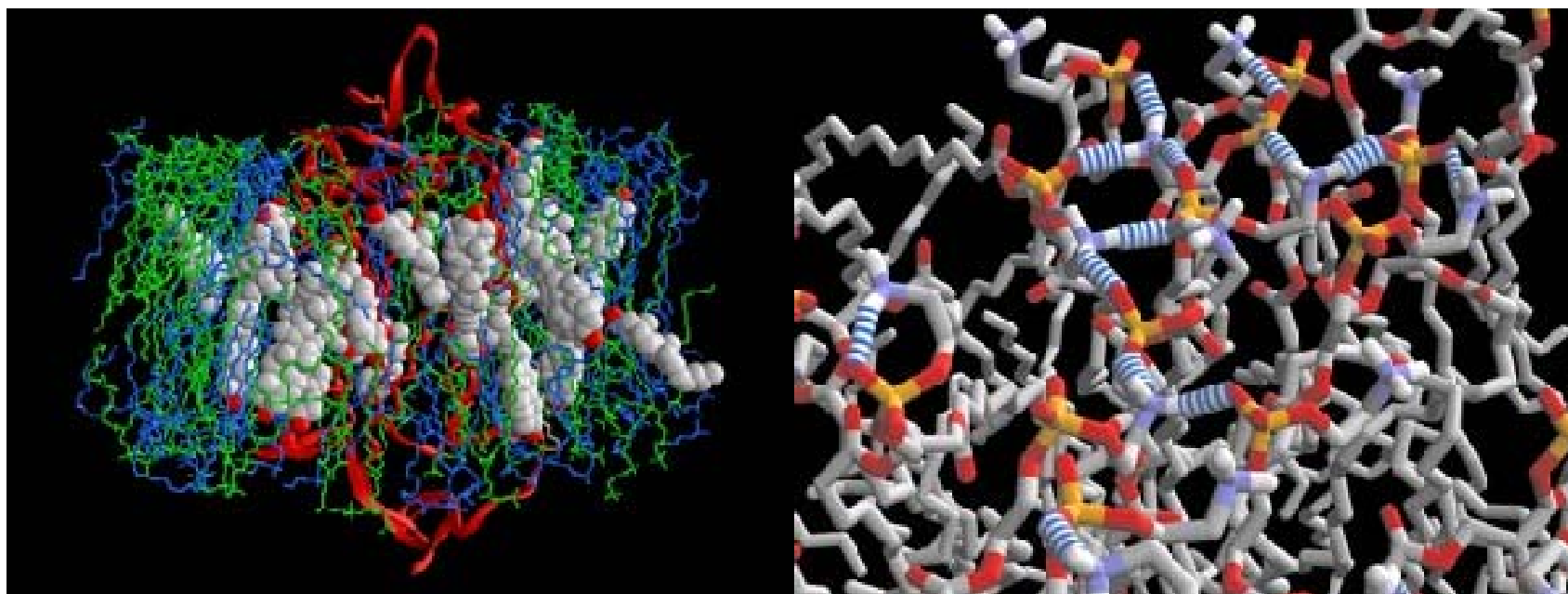






IBM's BlueGene/L: world's fastest supercomputer, 3 years running

596 TFLOPS (trillion floating point operations per second)



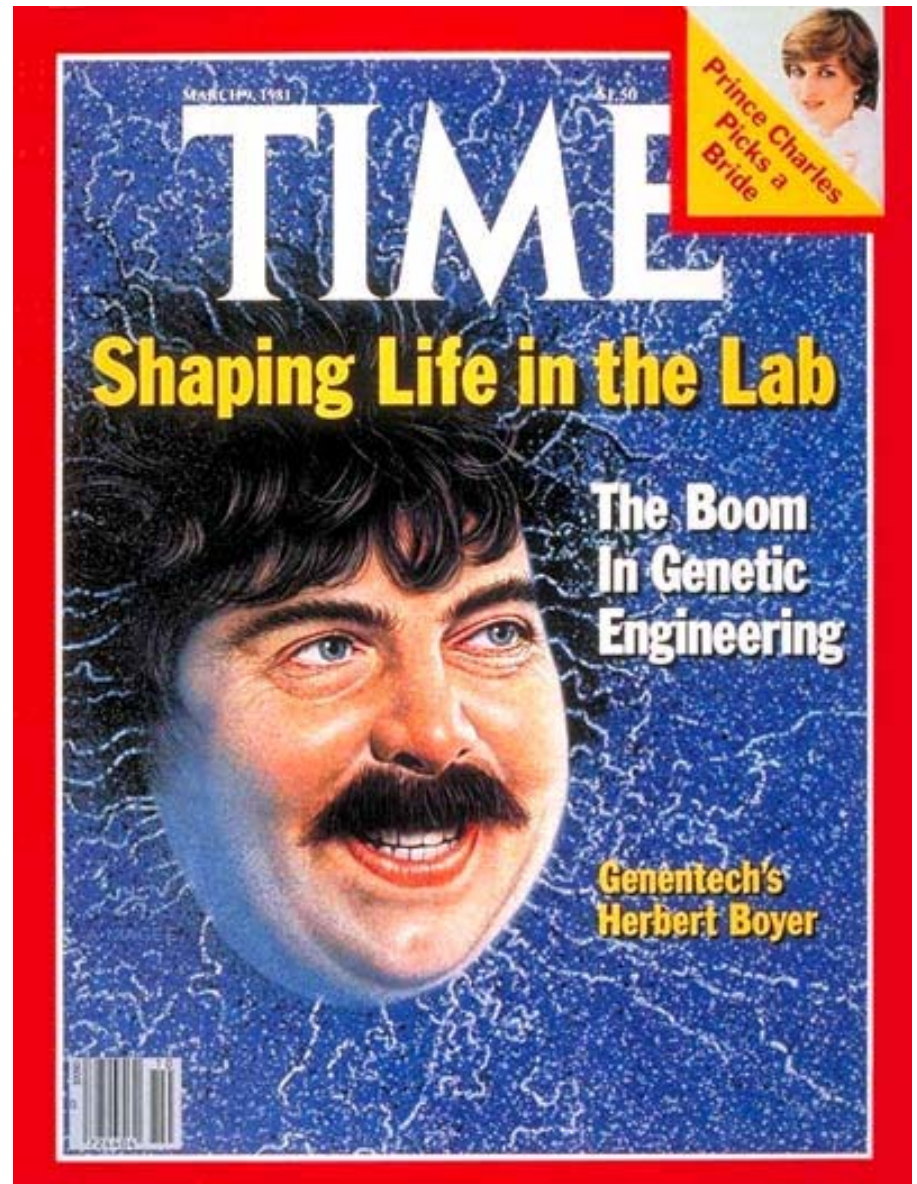
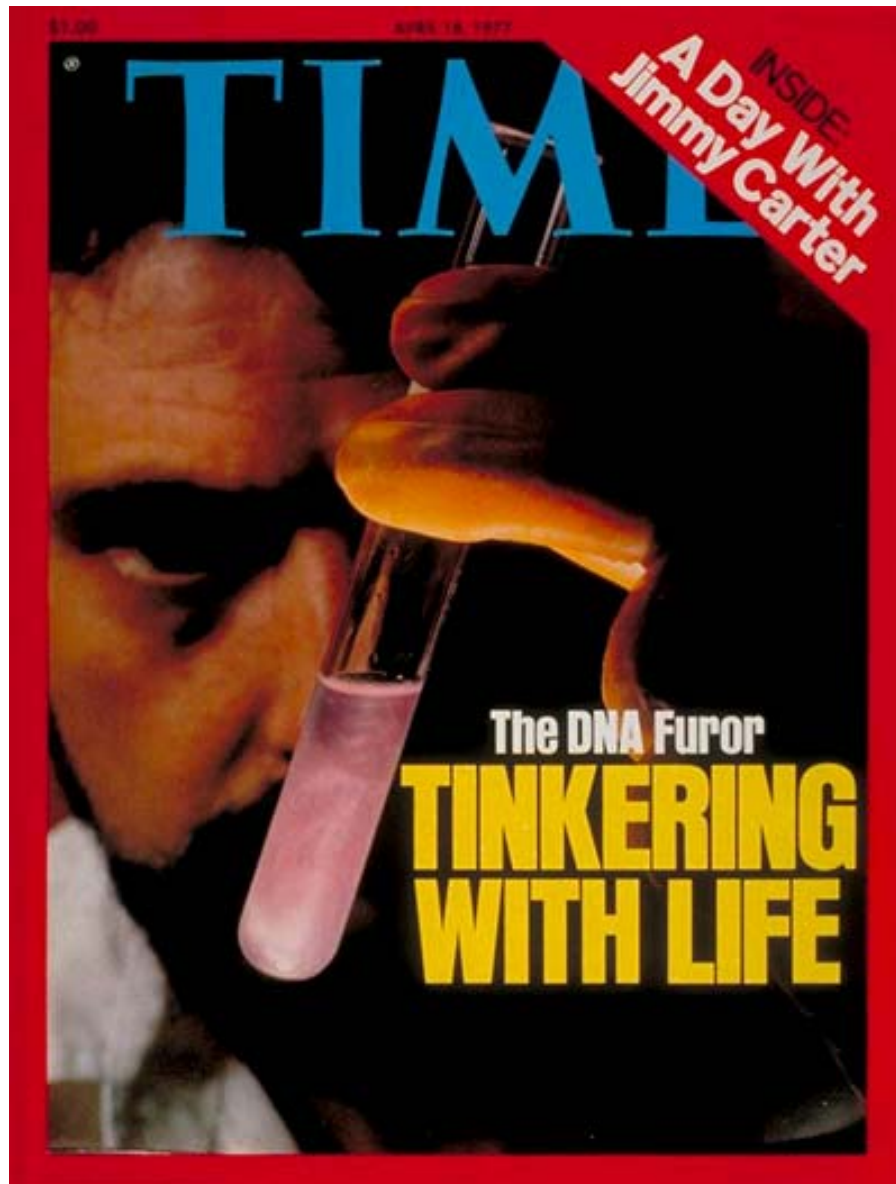
*The problem with reverse engineering...*

# REDUCTION | COMPLEXITY



*If we didn't build it, we may not be able to fully understand it.*

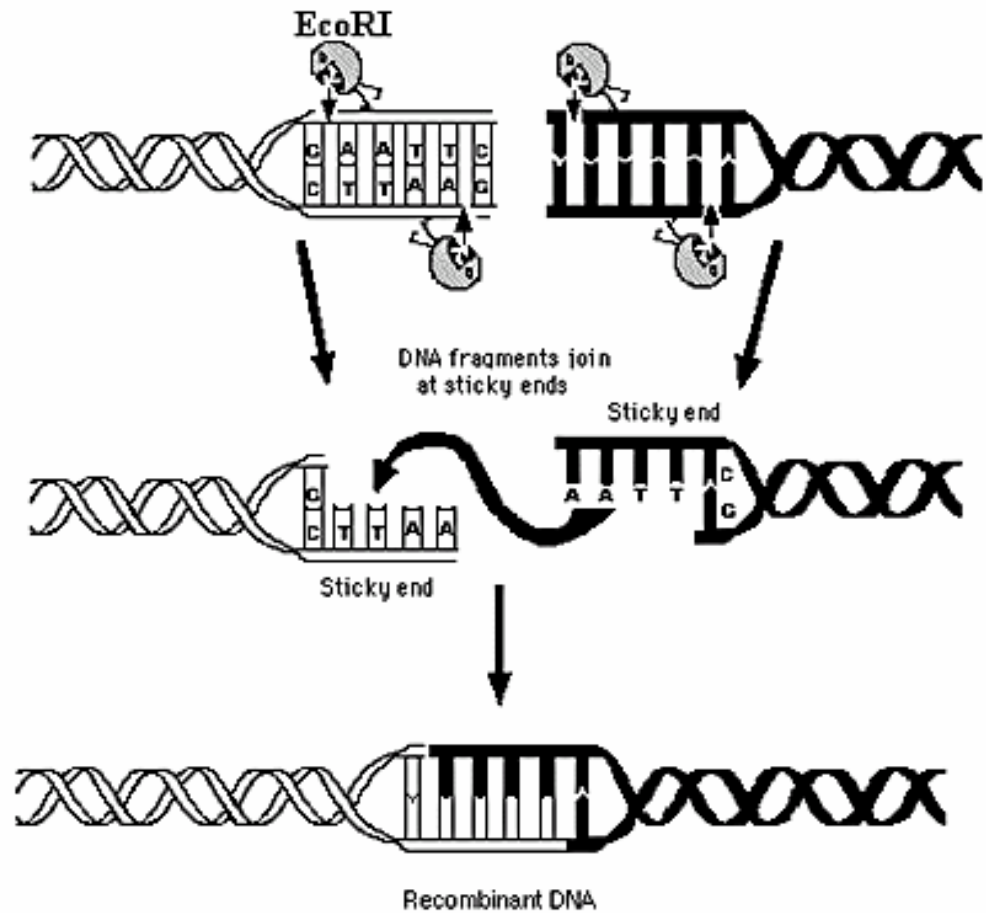
Towards “forward” bio-engineering







Over 3500 RE's available



**Restriction Enzyme  
Action of EcoRI**

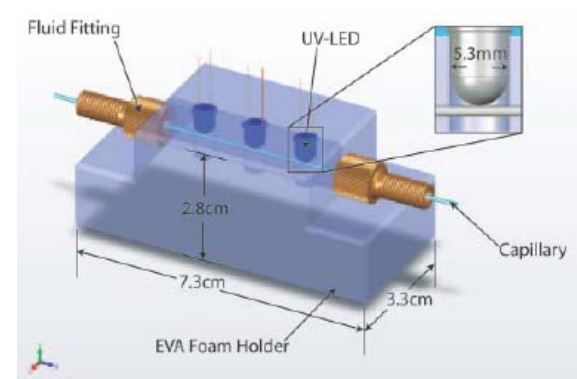
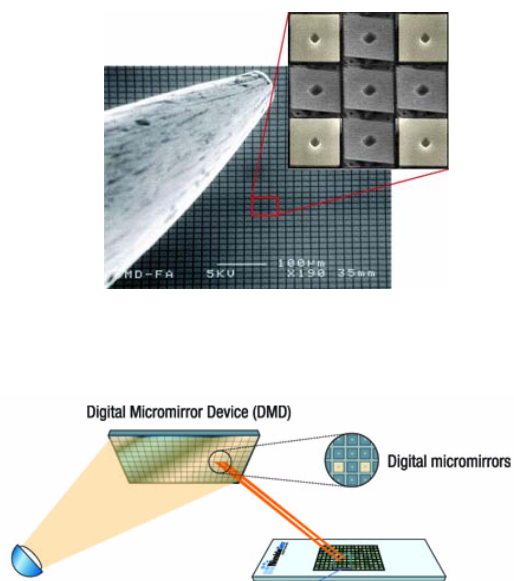
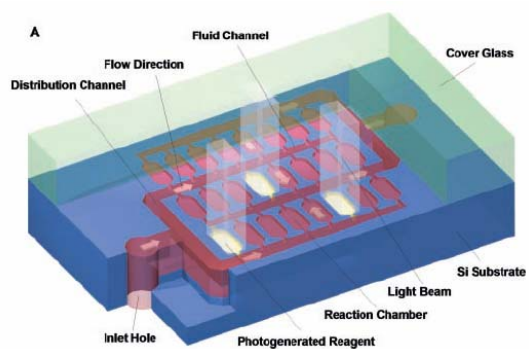


if you can **W**RiTe **D**Na,

You 'rE **n**O **LONGER** liMi**TED**

to "What **IS** " but To what you could MAKE •







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[www.blueheronbio.com](http://www.blueheronbio.com)

### **Bio S&T Gene Synthesis**

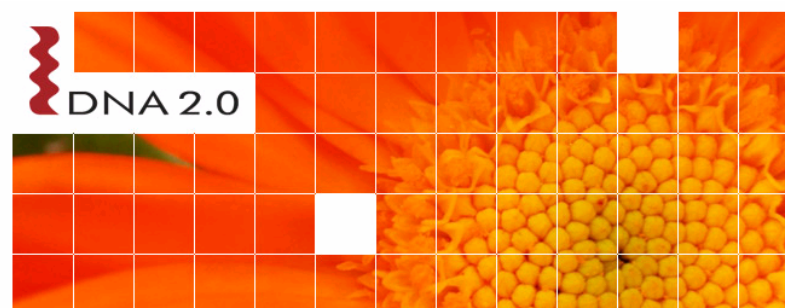
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and bonus DNA ladder.  
[www.biost.com](http://www.biost.com)

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### **Custom Gene Synthesis**

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to speed up R&D? Build with us.  
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THE BUILDING BLOCKS OF LIFE. BUILT FOR YOU.

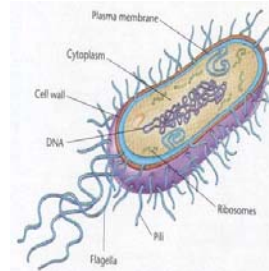
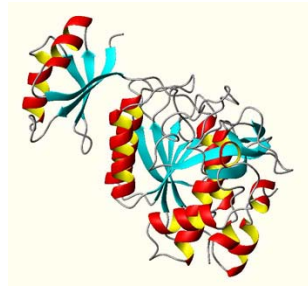


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# Applications dependent on synthetic capabilities and \$



single genes

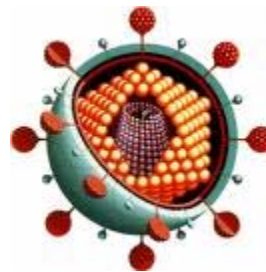
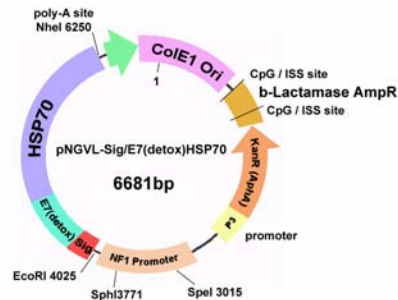
minimal life

base  
pairs



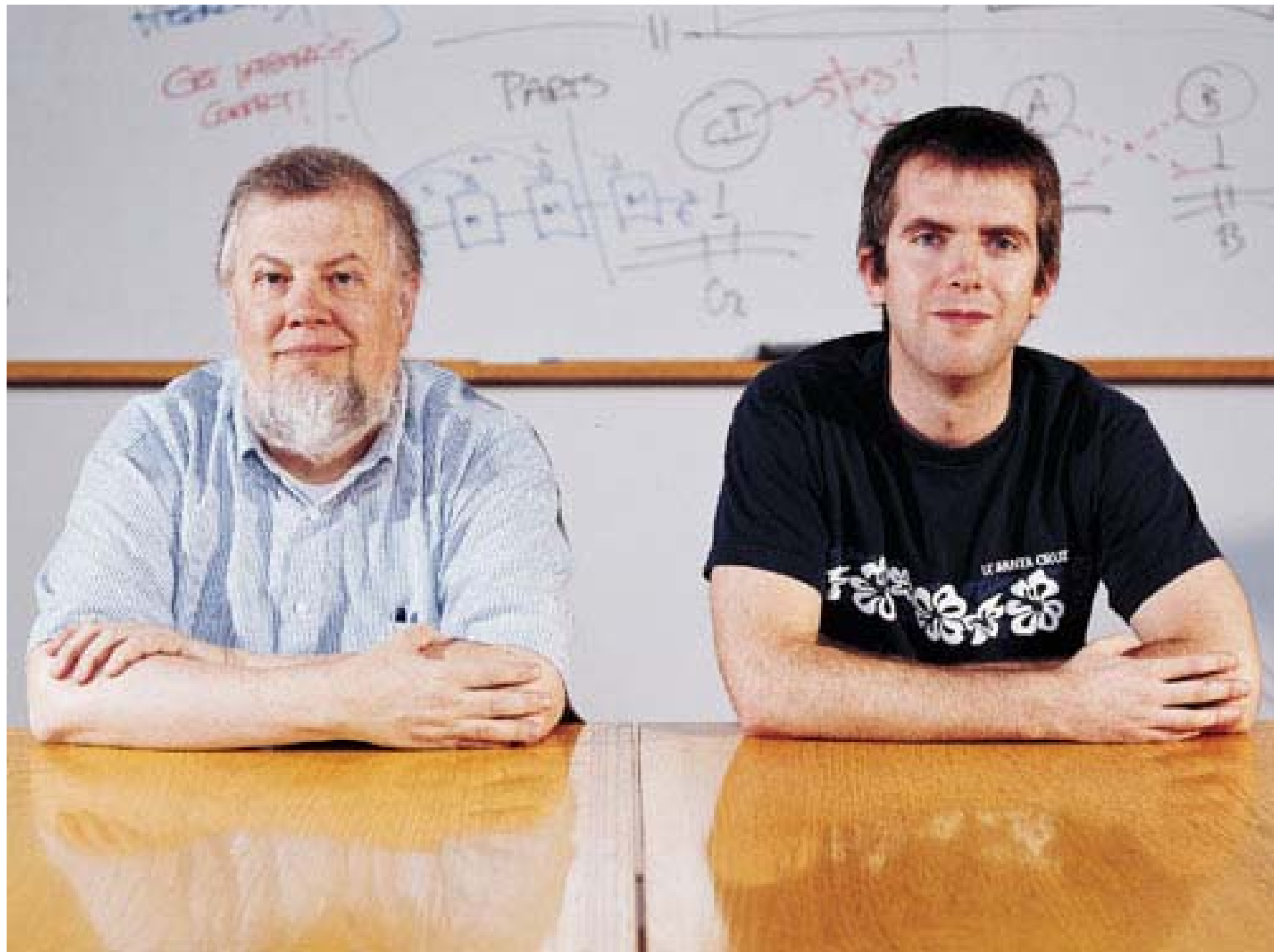
genetic circuits, viruses, GEMs

Engineered organisms
















## Systems

-  Measurement ?
  -  Measurement (Under Development) ?
  -  Projects(empty)
- 

## Devices

-  Reporters ?
  -  Protein Generator ?
  -  Inverters ?
  -  Composite Devices ?
  -  Signalling ?
  -  Measurement ?
- 

## Parts

-  Ribosome Binding Sites ?
-  Protein Coding ?
-  Regulatory ?
-  Terminators ?
-  RNA ?
-  Conjugation ?
-  DNA ?

## Chassis

-  E.coli Strains ?
-  Cell-Free Systems ?

## Mammalian

## Vectors

-  Plasmids ?
- 

## Other

-  Yeast Parts ?

## A.B Construction

### Intermediate ?

-  PCR Primer ?

-  Tags ?

-  Other

-  Deleted

-  Bacteriophage T7



jump to part

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## Transcriptional Regulators

### Available repressible regulators (normally ON) -?-

[Show 0 more parts](#)

[Edit](#)

-?-	Name	Description	Direction	Control -?-	Output Low High	Length
<a href="#">A</a> <a href="#">W</a>	<a href="#">BBa_I14032</a>	promoter P(Lac) IQ	Forward			37
<a href="#">A</a> <a href="#">W</a>	<a href="#">BBa_R0040</a>	promoter (tetR, negative)	Forward	aTc, tetracycline		54
<a href="#">A</a> <a href="#">W</a>	<a href="#">BBa_R0051</a>	promoter (lambda cl regulated)	Forward	lambda cl		49

### Available inducible regulators (normally OFF) -?-

[Show 0 more parts](#)

[Edit](#)

-?-	Name	Description	Direction	Control -?-	Output Low High	Length
<a href="#">A</a>	<a href="#">BBa_I12007</a>	Modified lambda Prm promoter (OR-3 obliterated)	Forward	cl		82
<a href="#">A</a>	<a href="#">BBa_R0062</a>	Promoter (luxR & HSL regulated -- lux pR)	Forward	luxR, HSL		55
<a href="#">A</a>	<a href="#">BBa_R0079</a>	Promoter (LasR & PAI regulated)	Forward	PAI		157
<a href="#">A</a>	<a href="#">BBa_R0080</a>	Promoter (AraC regulated)	Forward	araC		149

### Available other regulators

[Show 172 more parts](#)

[Edit](#)

-?-	Name	Description	Direction	Control -?-	Output Low High	Length
<a href="#">A</a> <a href="#">W</a>	<a href="#">BBa_I0500</a>	Inducible pBad/araC	Forward	araC, arabinose		1210
<a href="#">A</a> <a href="#">W</a>	<a href="#">BBa_I13453</a>	Pbad promoter				130
<a href="#">A</a> <a href="#">W</a>	<a href="#">BBa_J13002</a>	TetR repressed POPS/RIPS generator	Forward	ATc		74
<a href="#">A</a> <a href="#">W</a>	<a href="#">BBa_J13023</a>	3OC6HSL+LuxR dependent POPS/RIPS generator				117
<a href="#">A</a> <a href="#">W</a>	<a href="#">BBa_J23100</a>	constitutive promoter family member				35
<a href="#">A</a> <a href="#">W</a>	<a href="#">BBa_J23101</a>	constitutive promoter family member				35
<a href="#">A</a> <a href="#">W</a>	<a href="#">BBa_J23102</a>	constitutive promoter family member				35
<a href="#">A</a> <a href="#">W</a>	<a href="#">BBa_J23103</a>	constitutive promoter family member				35
<a href="#">A</a> <a href="#">W</a>	<a href="#">BBa_J23104</a>	constitutive promoter family member				35
<a href="#">A</a> <a href="#">W</a>	<a href="#">BBa_J23105</a>	constitutive promoter family member				35
<a href="#">A</a> <a href="#">W</a>	<a href="#">BBa_J23106</a>	constitutive promoter family member				35
<a href="#">A</a> <a href="#">W</a>	<a href="#">BBa_J23107</a>	constitutive promoter family member				35
<a href="#">A</a> <a href="#">W</a>	<a href="#">BBa_J23108</a>	constitutive promoter family member				35
<a href="#">A</a> <a href="#">W</a>	<a href="#">BBa_J23109</a>	constitutive promoter family member				35
<a href="#">A</a> <a href="#">W</a>	<a href="#">BBa_J23110</a>	constitutive promoter family member				35
<a href="#">A</a> <a href="#">W</a>	<a href="#">BBa_J23111</a>	constitutive promoter family member				35

<http://parts.mit.edu>

# BBa\_F2620

3OC<sub>6</sub>HSL → PoPS Receiver

[http://parts.mit.edu/registry/index.php/Part:BBa\\_F2620](http://parts.mit.edu/registry/index.php/Part:BBa_F2620)



Authors:  
Barry Cantor ([bcantor@mit.edu](mailto:bcantor@mit.edu))  
Anna Labno ([labnoa@mit.edu](mailto:labnoa@mit.edu))

Last Update: 15 January 2007

## Description

A transcription factor (LuxR, BBa\_C0062) that is active in the presence of cell-cell signaling molecule 3OC<sub>6</sub>HSL is controlled by a TetR-regulated operator (BBa\_R0040). Device Input is 3OC<sub>6</sub>HSL. Device output is PoPS from a LuxR-regulated operator. If used in a cell containing TetR then a second input signal such as aTc can be used to produce a Boolean AND function.

## Characteristics

Input Swing: 1E-9 to 1E-6 M 3OC<sub>6</sub>HSL, exogenous

Output Swing: 0±1 to 503±1 GFP molecules cfr<sup>-1</sup> s<sup>-1</sup>

Switch Point: 7±1 nM 3OC<sub>6</sub>HSL, exogenous

LH Response: 9 min (t<sub>50%</sub>), 27 min (t<sub>90%</sub>)

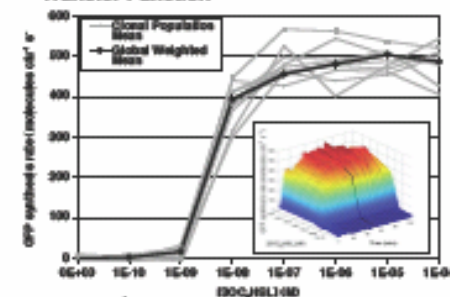
## Key Parts

BBa\_R0040: TetR-regulated operator

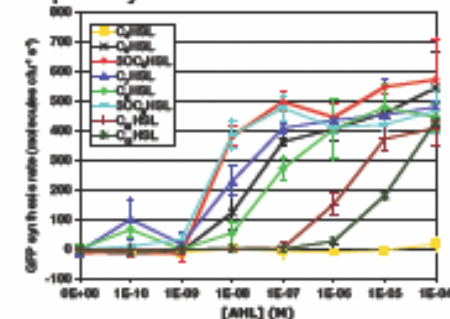
BBa\_C0062: luxR ORF

BBa\_R0002: LuxR-regulated operator

## Transfer Function\*



## Specificity\*



## Demand (low/high Input)

Translational: 256/8048 ribosomes cfr<sup>-1</sup>  
3.8E3/1.2E5 charged tRNA cfr<sup>-1</sup> s<sup>-1</sup>

## Compatibility

Chassis: Compatible with MC4100, MG1655, and DH5α

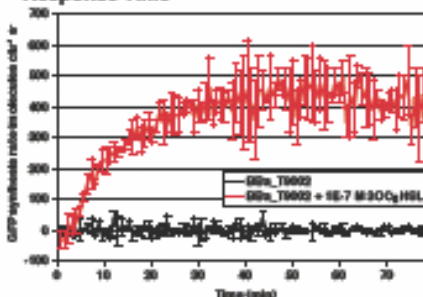
Plasmids: Compatible with pSB3K3 and pSB1A2

Devices: Compatible with E0240, E0430 and E0434

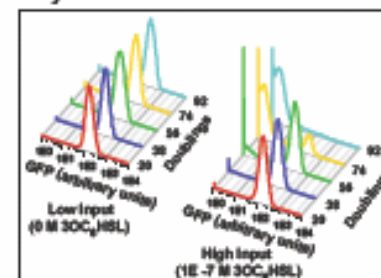
Crosstalk with systems containing TetR (C0040)

Signaling: Crosstalk with input molecules similar to 3OC<sub>6</sub>HSL

## Response Time\*



## Stability\*\*



## Stability (low/high Input)

Genetic: >32/74 replication events\*\*

Performance: >32/74 replication events\*\*

## Conditions (abridged)

Output: Indirect via BBa\_E0240

Vector: pSB3K3

Chassis: MG1655

Culture: Supplemented M9, 37°C

\*Equipment: PE Victor3 plate reader

\*\*Equipment: BD FACScan cytometer

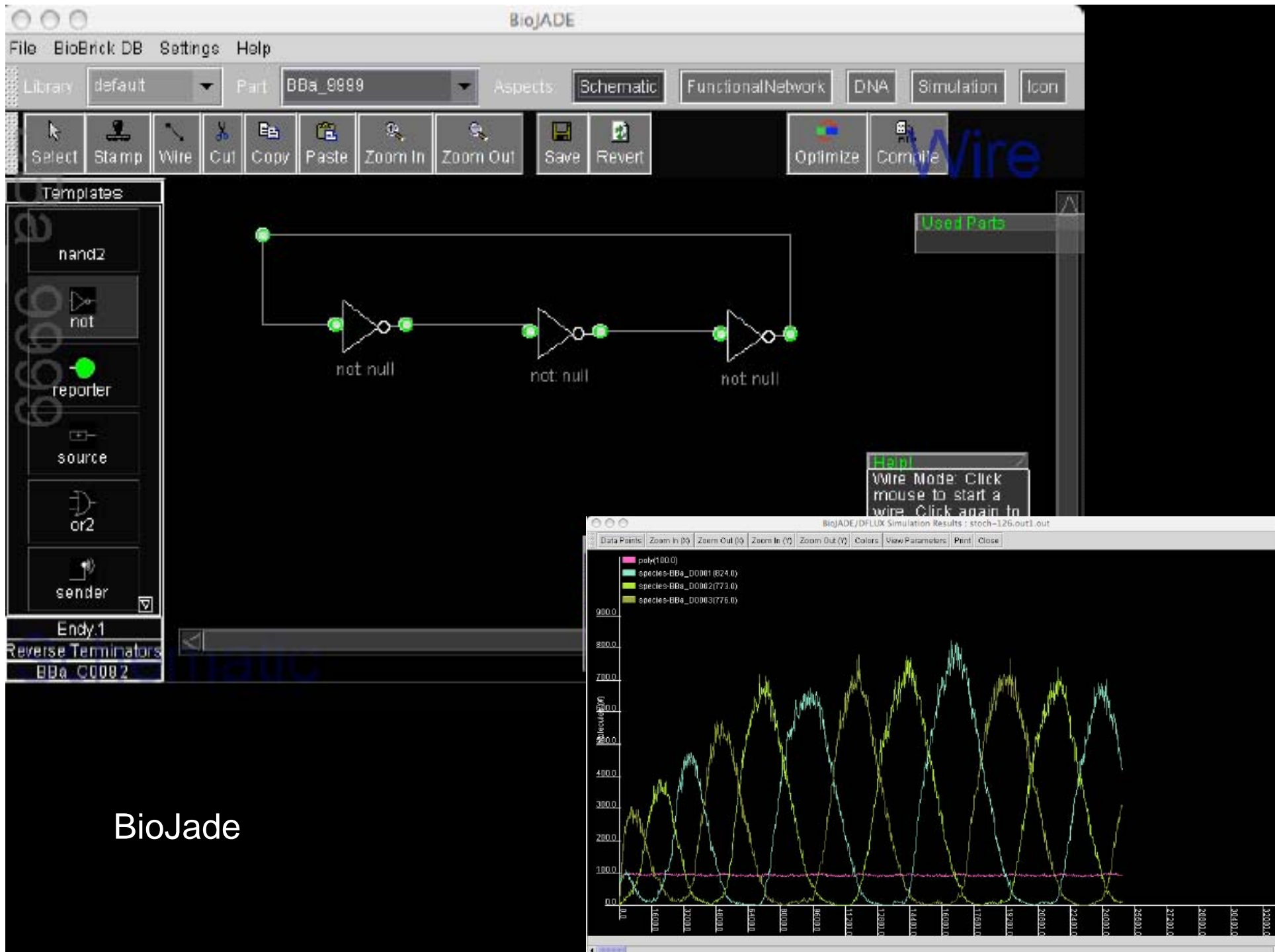
Signaling Devices

Registry of Standard Biological Parts

making life better, one part at a time

License: Public



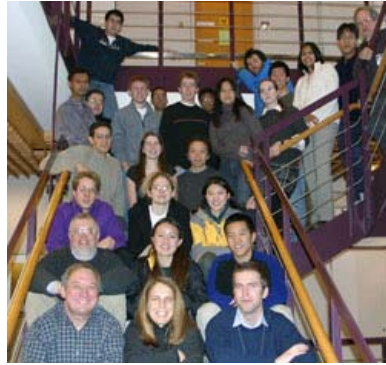


BioJade

*Synthetic biology requires an unprecedented level of interdisciplinary cooperation, fewer laboratory skills, and less overhead than conventional genetic engineering.*













# iGEM 2007 Wiki

International Genetically Engineered Machine Competition

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## Cambridge & Melbourne Leap Together

Cambridge University & Melbourne University leap together after co-winning the Best BioBrick Part award.

## iGEM?

Hundreds of undergraduates all over the world spend their summer making Synthetic Biology a reality by participating in the annual International Genetically Engineered Machine competition.

iGEM through the years

- 2008
- 2007
- 2006

[Learn More](#)

## Results of the Jamboree

sat & sun, nov 3-4



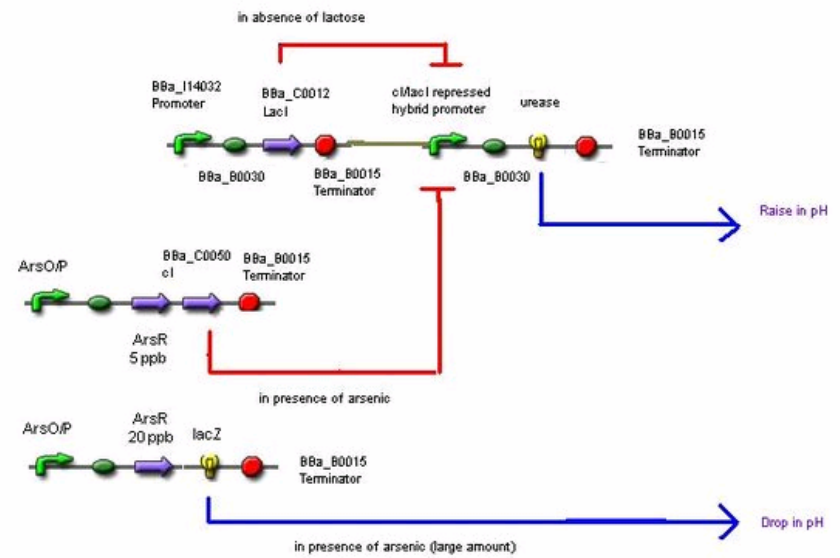
iGEM 2007 is now  
officially concluded!  
Congratulations to all!

- [Results](#)
- [See the medal winners](#)
- [Media](#) (including links to videos and [flickr™](#) photo gallery)
- [Learn about iGEM 2008](#)

## calendar

Jamboree roster + fees due	<i>fri</i> 12 oct 07
iGEM wiki frozen + parts postmarked	<i>fri</i> 26 Oct 07
Jamboree!	<i>sat-sun</i> 3-4 nov 07
Registry + BioBrick foundation workshops	<i>sun-tue</i> 4-6 nov 07

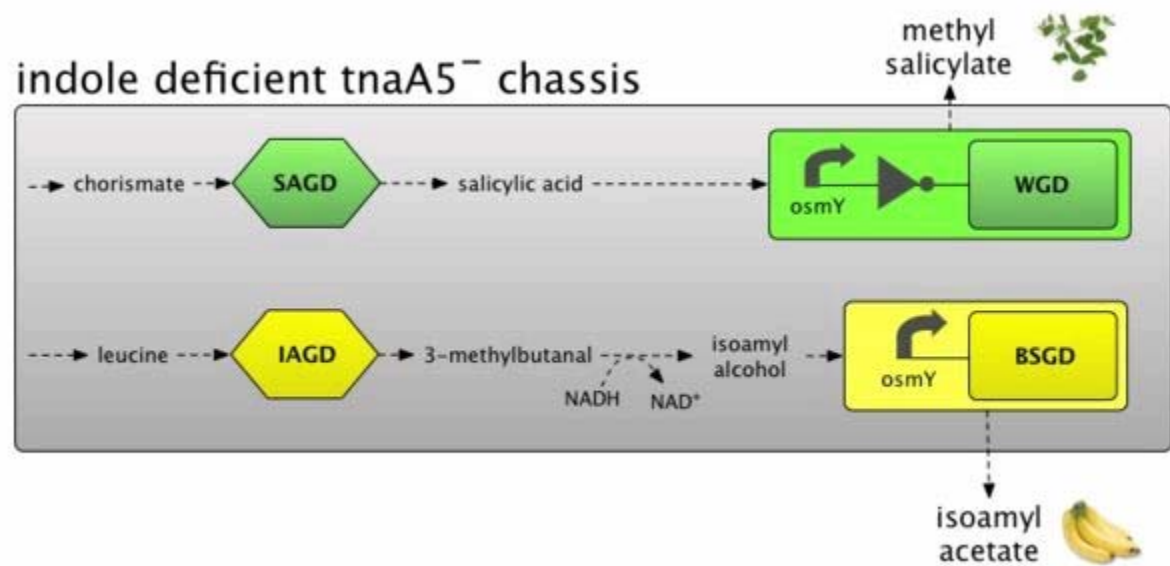








# indole deficient $tnaA5^-$ chassis





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### Sci-Tech

Under a magnification of 6836x, this scanning electron micrograph (SEM) depicts a number of Gram-negative *Escherichia coli* bacteria of the strain O157:H7. (Courtesy of Centers of Disease Control and Prevention)

**Alberta team trying to turn E. coli into fuel**

Updated Mon. Nov. 5 2007 8:12 AM ET

*The Canadian Press*

CAMBRIDGE, Mass. — A University of Alberta team trying to turn E. coli into fuel has earned a first place finish in the energy category at an international genetic engineering competition at MIT.

Andrew Hessel, a consultant with the Alberta Ingenuity Fund who has helped teams from three Alberta universities compete, says while none of the teams from the University of Calgary, the University of Lethbridge or the U of A made it to the final competition, he calls it a fantastic educational experience.

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**USER TOOLS**

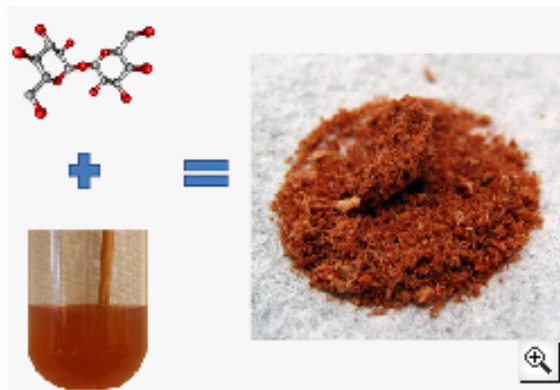
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## Genetic-Engineering Competitors Create Modular DNA Dev Kit

By Alexis Madrigal  11.13.07 | 7:30 PM



University of California-Berkeley students created "bacto-blood," which can be freeze-dried and may one day be used for blood transfusions.

*Image: Austin Day*

College and high school students are helping MIT scientists develop an open source development kit for biological systems that could do for cells what Linux has done for computers.

As part of the International Genetically Engineered Machines competition held in Cambridge, Massachusetts, last week, Peking University students created tiny assembly lines out of bacteria. Their entry, "[Towards a Self-Differentiated Bacterial Assembly Line](#)," won them the grand prize among 50 teams from around the world.

"Biology is going to be able to make the things that we want," said Tom Knight, an MIT engineer and co-founder of iGEM. "And when that happens, the economics of production are going to change dramatically. It doesn't

take a billion-dollar [facility] to make stuff. It takes a hundred-dollar incubator."



# The New York Times

English, Algebra, Phys Ed ... and Biotech



Jim Wilson/The New York Times

George Cachianes, left, formerly of Genentech, teaches biotechnology at Lincoln High School in San Francisco.

# BERKELEY CENTER FOR SYNTHETIC BIOLOGY

A JOINT PROGRAM OF THE CALIFORNIA INSTITUTE FOR QUANTITATIVE BIOMEDICAL RESEARCH (QB3)  
AND LAWRENCE BERKELEY NATIONAL LABORATORY (LBNL)

The California Institute for Quantitative Biomedical Research (QB3) and Lawrence Berkeley National Laboratory (LBNL) have joined forces to accelerate the growth of synthetic biology, a new field that promises major new advances in preventing and treating disease, generating new energy sources, and preventing and mitigating environmental threats.

Opening in spring 2005 in a spacious, modern building in west Berkeley, the Berkeley Center for Synthetic Biology gives renowned scientists and engineers the chance to pool their talents and collaborate in new ways, with enormous potential benefits for California's citizens in the form of advances in biomedicine and energy renewables and economic growth.

Synthetic biologists study the control and design of biological components and new organisms to solve a host of important health, energy, and environmental problems that cannot be solved using naturally occurring biological entities. The inherently



QB3 and LBNL scientists occupy lab space in a building renovated in 1997 for biotech research, previously leased by Bayer, featuring large labs, viral suites, and tissue culture rooms. UCSF Mission Bay and numerous biotech firms are nearby.



## MIT establishes groundbreaking biological engineering major

February 17, 2005

The Massachusetts Institute of Technology faculty yesterday approved a new course of study for undergraduates, in biological engineering, the first entirely new curriculum established at the Institute in 29 years.

University of  
Lethbridge



MAY 3, 1982

\$1.50

# TIME

## COMPUTER GENERATION

A New Breed of Whiz Kids



**-3%**  
INFLATION VANISHES!  
At Least for  
A Month



**TRS-80 COMPUTER** CAT. NO. 68-2030

# Whizkids™

ALEC AND SHANNA STARRING IN

## THE COMPUTER TRAP

**COMPLIMENTS OF Radio Shack**  
The Name in Classroom Computing

DICK AYERS AND CHIC STONE

THAT'S RIGHT, ALEC! SCRIPTSIT IS A WORD PROCESSING PROGRAM. MY DAD HAS A TRS-80 MODEL 12 COMPUTER WITH SCRIPTSIT IN HIS OFFICE... AND HE TAUGHT ME HOW TO USE IT WITH A DAISY WHEEL PRINTER...

...TO WRITE BUSINESS LETTERS, RESEARCH NOTES, PRESS RELEASES, AND BULLETINS.

SHANNA YOU KNOW SO MUCH - SHOW US HOW...

...SCRIPTSIT WORD PROCESSING WORKS IN OUR SCHOOL'S OFFICE.

**IN THE SCHOOL OFFICE...**

TURN ON THE POWER SWITCH THEN "INSERT DISKETTE"... CAREFULLY PUSH DISKETTE INTO THE SLOT (DRIVE 0) AND ROTATE THE LATCH TO A HORIZONTAL POSITION.

DRIVE 0  
DRIVE 1

AFTER THAT THE WORD "INITIALIZING" APPEARS WHICH MEANS THE COMPUTER IS LOADING THE PROGRAM...

INITIALIZING

... ALSO, THE SMALL RED LIGHT NEXT TO THE DISK DOOR WILL BE "ON"!

AFTER THE LIGHT GOES OUT, THE PROGRAM HAS BEEN "LOADED" INTO THE COMPUTER. NEXT, THE COMPUTER TELLS YOU TO TYPE IN THE DATE...

... FOR EXAMPLE APRIL 6, 1984, TYPE 04/06/1984 AND THEN PRESS THE [ENTER] KEY.

ENTER DATE (MM/DD/YYYY)

NEXT, THE COMPUTER PROMPTS YOU TO ENTER THE TIME USING THE 24-HOUR SYSTEM, GIVING HOURS, MINUTES AND SECONDS.

FOR EXAMPLE 9:30 AND 20 SECONDS A.M., TYPE THIS WAY-- 09.30.20. AND THEN PRESS THE [ENTER] KEY.

ENTER TIME (HH. MM. SS)

THAT MAKES THE "DIRECTORY" APPEAR ON THE SCREEN. THE DIRECTORY IS DIVIDED INTO SIX "CELLS". EACH CELL IS THE STORAGE UNIT FOR INFORMATION ABOUT ONE DOCUMENT...

NAME	DATE	TIME	STATUS	LENGTH	BYTES	DISK	FILE
SCRIPTSIT	04/06/1984	09:30:20	OK	1000	1000	1	1
SCRIPTSIT	04/06/1984	09:30:20	OK	1000	1000	1	1
SCRIPTSIT	04/06/1984	09:30:20	OK	1000	1000	1	1
SCRIPTSIT	04/06/1984	09:30:20	OK	1000	1000	1	1
SCRIPTSIT	04/06/1984	09:30:20	OK	1000	1000	1	1
SCRIPTSIT	04/06/1984	09:30:20	OK	1000	1000	1	1

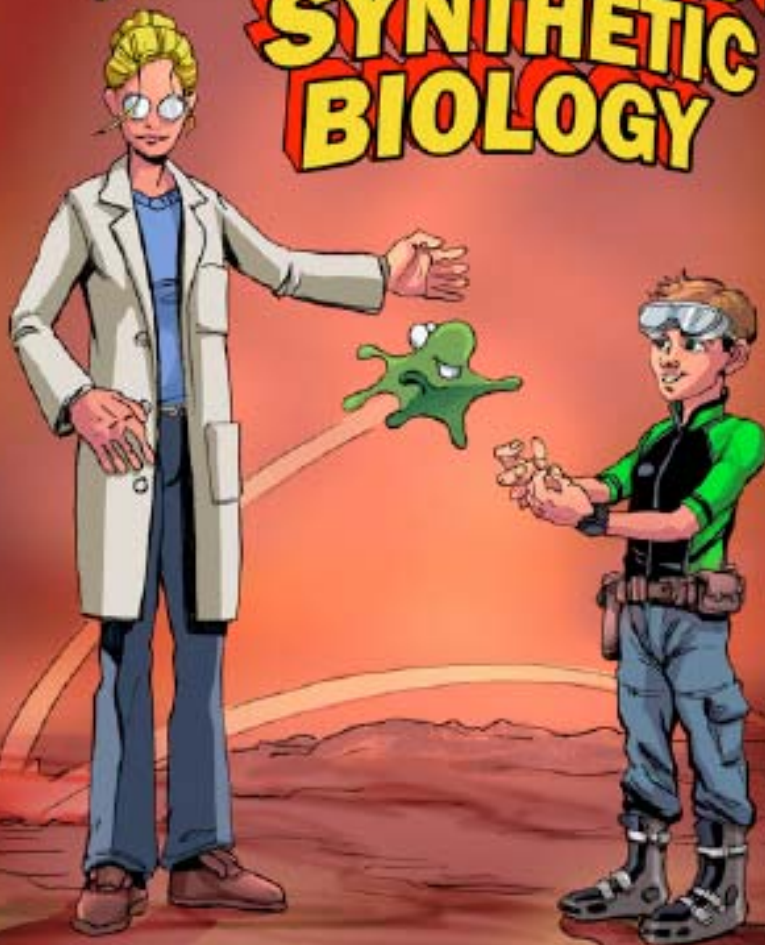
... AND IN TURN, A DOCUMENT CAN BE MADE UP OF SEVERAL PAGES OF INFORMATION.

IS THERE A SCRIPTSIT PROGRAM FOR OUR CLASSROOM TRS-80 MODEL 4'S?

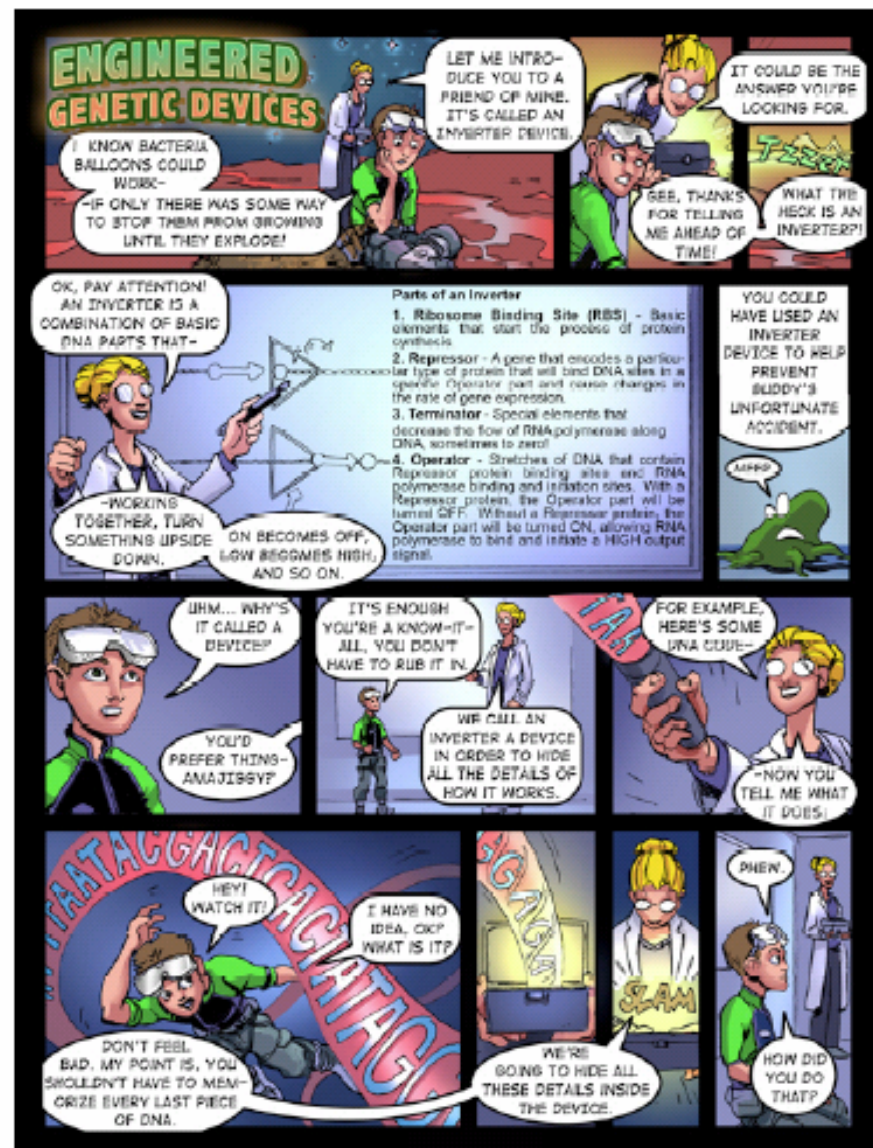
YES, THERE IS A SPECIAL SCRIPTSIT PROGRAM FOR THE MODEL 4'S!

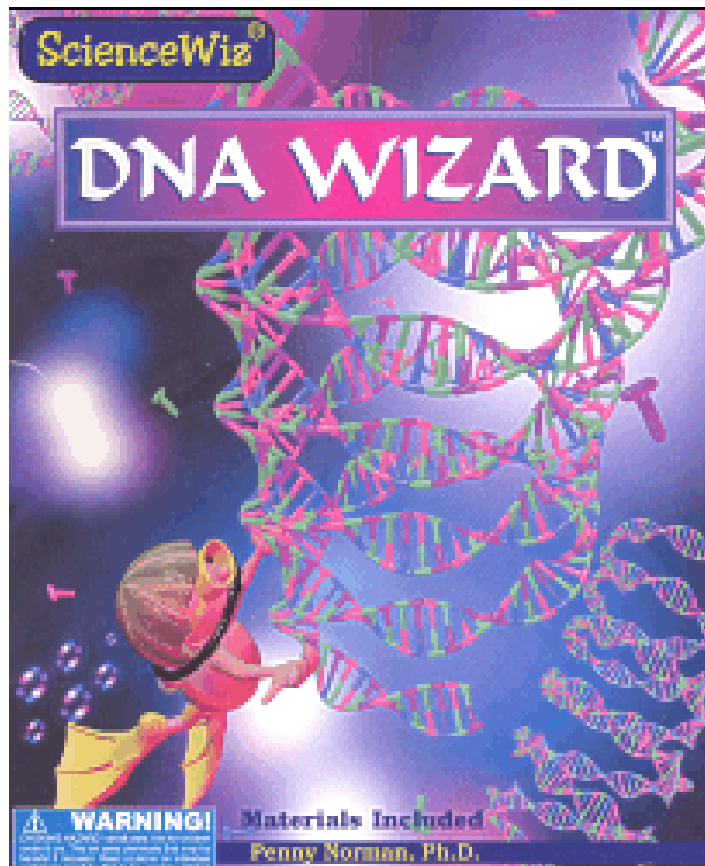


# ADVENTURES IN SYNTHETIC BIOLOGY



STORY: DREW ENDY ISADORA DEESE  
THE MIT SYNTHETIC BIOLOGY WORKING GROUP  
ART: CHUCK WADEY [WWW.CHUCKWADEY.COM](http://WWW.CHUCKWADEY.COM)





# Projects with DNA

**For ages 8 and up  
Adult Supervision Required**

Materials included except for the items listed.  
Through play, hands-on projects, patterns and puzzles  
this book and kit explores the amazing DNA story.

**Extract DNA**

**Heat SHOCK!**

**Decode the code of life**

**Build a DNA ladder.**

**Grow glowing cells**

**Is it a boy or girl?**

**Solve the chromosome puzzle.**

**Ooey, Gooley, DNA!**

**Dress up for sterile techniques.**

**Quality time, quality learning, quality play.**



# Why do bioengineering?

- Any chemical process can be done with biochemistry – no need for harsh chemical conditions, temperature, etc.
- Runs on sugar or sunlight
- Bio-compatible and biodegradable
- Self-reproducing > very low cost
- Potentials are virtually untapped
- Evidence suggests many of today's engineering activities will begin to move towards bioengineered processes > big economic shift on the order of computing
- Life and life processes are important – should be widely used and understood, not in the hands of a few elites – hence the need for “open source biology”



## Alberta Ingenuity operates

the Alberta Heritage Foundation for Science and Engineering Research,  
a \$1 billion endowment to support research.

**Ingenuity launches bold new \$100 million research program**

October 10, 2007 - Ingenuity Accelerators focus on ensuring future prosperity of the province.

- Registration fees (\$1000/team)
- Teach the teachers meeting (MIT, May)
- Regional meetings and seminars
- Travel and accommodation for up to 100 students to MIT Jamboree in November
- \$5,000 research start-up fees
- Support, technical advice, media relations